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# Effect of reproduction on escape responses, metabolic rates and muscle mitochondrial properties in the scallop *Placopecten magellanicus*

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#### Abstract:

In scallops, gametogenesis and spawning can diminish the metabolic capacities of the adductor muscle and reduce escape response performance. To evaluate potential mechanisms underlying this compromise between reproductive investment and escape response, we examined the impact of reproductive stage (pre-spawned, spawned and reproductive quiescent) of the giant scallop, Placopecten magellanicus, on behavioural (i.e., escape responses), physiological (i.e., standard metabolic rates and metabolic rates after complete fatigue) and mitochondrial capacities (i.e., oxidative rates) and composition. Escape responses changed markedly with reproductive investment, with spawned scallops making fewer claps and having shorter responses than pre-spawned or reproductive-guiescent animals. After recuperation, spawned scallops also recovered a lower proportion of their initial escape response. Scallop metabolic rate after complete fatigue (VO<sub>2max</sub>) did not vary significantly with reproductive stage whereas standard metabolic rate (VO<sub>2min</sub>) was higher in spawned scallops. Thus spawned scallops had the highest maintenance requirements (VO<sub>2min</sub>/VO<sub>2max</sub>). Maximal capacities for glutamate oxidation by muscle mitochondria did not change with reproductive stage although levels of ANT and cytochromes as well as cytochrome C oxidase (CCO) activity did. Total mitochondrial phospholipids, sterols and the proportion of phospholipid classes differed only slightly between reproductive stages. Few modifications were detected in the fatty acid (FA) composition of the phospholipid classes except in cardiolipin (CL). In this class, prespawned and spawned scallops had fairly high proportions of 20:5n-3 whereas this FA in reproductivequiescent scallops was threefold lower and 22:6n-3 was significantly higher. These changes paralleled the increases in CCO activity and suggest an important role of CL on the modifications of CCO activity in scallops. However, mitochondrial properties could not explain the decreased recuperation ability from exhausting exercise in spawned scallops. Shifts in maintenance requirements (VO<sub>2min</sub>/VO<sub>2max</sub>) and aerobic scope (VO<sub>2max</sub> - VO<sub>2min</sub>) provided the best explanation for the impact of reproduction on escape response performance.

**Keywords:** mitochondria, reproduction, scallops, muscle, escape response, metabolic rate, cytochrome C oxidase, phospholipids, plasmalogens, fatty acids, sterols, *Placopecten magellanicus*.

#### 27 Introduction

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29 The requirements of broadcast spawning lead many marine invertebrates to invest so 30 heavily in gametogenesis and spawning that reproduction represents a major stress (Barber 31 and Blake 1991). Given the material and energetic costs of gametogenesis and spawning, 32 metabolic expenditures are likely to change considerably during the reproductive cycle, 33 particularly in species with a high reproductive output. In scallops, gonadal maturation and 34 spawning lead to a negative energetic balance, a generally diminished physiological condition 35 and a decreased capacity to recover from exhausting exercise (Barber and Blake 1985; 36 Brokordt et al. 2000a; Brokordt et al. 2000b). Thus, scallops face a compromise between two 37 major components of their fitness: reproductive success and individual survival.

38 In scallops, the striated adductor muscle is one of the largest soft tissues and has as its 39 primary role the rapid movement of valves during escape responses. The capacities of the 40 adductor muscle are thus important for survival during encounters with predators. In many 41 scallop species, the energetic reserves of the adductor muscle are depleted during gonadal 42 maturation and spawning (Barber and Blake 1981; Barber and Blake 1991; Brokordt and 43 Guderley 2004). At the same time, the glycolytic and aerobic capacity of the adductor muscle 44 declines, which could explain reductions in the scallops' swimming ability, and particularly in 45 their capacity to recuperate from exhaustive exercise (Brokordt et al. 2000a; Brokordt et al. 46 2000b; Brokordt and Guderley 2004). The impact of the reproductive cycle upon escape 47 response performance could also reflect the instantaneous metabolic costs of reproductive 48 investment. Indeed, the gametogenetic cycle of scallops accentuates seasonal changes in 49 metabolic rate with the highest rates occurring during reproductive months and the lowest 50 rates during winter quiescence (Shumway et al. 1988). Thus, the metabolic costs of 51 reproductive investment and spawning could reduce aerobic scope and slow aerobic52 recuperation.

53 Rapid valve movements (claps) in scallops are primarily fueled by phosphoarginine, 54 with subsequent recovery using anaerobic glycolysis and then oxidative metabolism. In 55 Placopecten magellanicus and Argopecten irradians concentricus, 70% of the ATP required 56 for exhaustive escape activity is derived from phosphoarginine, with only 30% arising from 57 anaerobic glycolysis and octopine generation (Thompson et al. 1980; de Zwaan et al. 1980; 58 Livingstone et al. 1981; Chih and Ellington 1986). Phosphoarginine levels decrease as a 59 function of the number of claps (Livingstone et al. 1981; Bailey et al. 2003). The complete 60 restoration of phosphoarginine pools after exhaustiing swimming requires aerobic metabolism 61 (Livingstone et al. 1981), presumably relying upon ATP production by adductor muscle 62 mitochondria (Guderley et al. 1995). Both muscle glycolytic capacities and the oxidative 63 capacities of muscle mitochondria change with the reproductive cycle in Euvola ziczac and 64 Chlamys islandica, decreasing in animals that have invested extensively in reproduction 65 (Boadas et al. 1997; Brokordt et al. 2000a). Thus, muscle mitochondria can be considered as 66 key organelles in covering the high energy demand of aerobic recuperation in scallops. Any 67 limitation of their oxidative capacity could hinder aerobic metabolism and could explain the 68 impact of reproductive investment on aerobic recuperation.

Mitochondrial capacities can be changed by shifts in their protein and phospholipid components. Alterations in phospholipid head groups and acyl chain compositions modify the molecular activities of respiratory chain components (Hazel 1972a; Hazel 1972b). Mitochondrial membrane proteins can have specific requirements for phospholipid head group arrangements and FA in their proximity (Clandinin et al. 1985; Berger et al. 1993; Schlame et al. 2000). Thus, modifications in membrane lipid composition, even of minor

phospholipid classes during the reproductive cycle, could underlie changes in mitochondrial
oxidative capacity with the reproductive cycle.

77 This study examined potential mechanisms underlying the compromise between 78 locomotor performance and reproductive investment in the giant scallop, P. magellanicus, by 79 evaluating changes in the aerobic power budget and mitochondrial capacities with 80 reproductive investment. To this end, we compared scallops sampled in their natural habitat 81 at 3 reproductive stages (pre-spawned, spawned and reproductive-quiescent). We 82 characterized their escape responses, recuperation from exhaustion, aerobic power budget, 83 muscle mitochondrial capacities and mitochondrial composition. To evaluate how respiratory 84 capacity could be associated with swimming ability, particularly in the capacity to recuperate from exhausting exercise, oxygen uptake was measured at rest in non-feeding animals 85 86  $(VO_2min)$  and during aerobic recovery from exhausting exercise  $(VO_2max)$ . We further 87 examined properties of adductor muscle mitochondria as possible drivers of whole animal 88 metabolic rate and recuperation from exhausting exercise. We determined maximal rates of 89 glutamate oxidation, levels of adenine nucleotide translocase (ANT), concentrations of 90 cytochromes A, B, C and C<sub>1</sub>, activity of cytochrome C oxidase (CCO) and proportions of the 91 phospholipid classes and subclasses as well as their FA compositions. By examining 92 performance and structure at these levels, we sought to evaluate whether changes in scallop 93 escape response performance with reproductive investment were due to their aerobic power 94 budget or to their muscle mitochondrial properties.

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#### 97 Materials and Methods

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# Collection and maintenance of experimental animals

100 The giant scallop, *Placopecten magellanicus*, population we studied is located at the 101 mouth of Baie des Chaleurs near Percé, Québec, in eastern Canada's Gulf of St. Lawrence, (48° 30' N, 65° 15' W). As at this location, spawning events occur between mid-July and 102 early September (Bonardelli et al. 1996), we sampled at three dates: July 5<sup>th</sup> (before spawning 103 events). September 6<sup>th</sup> (during or just after or spawning) and October 18<sup>th</sup> 2004 (reproductive 104 105 quiescence). At each sampling date, 9 scallops (shell height: 106.9±10.9 mm) were collected 106 by SCUBA diving at 30 m and water temperature was noted. Animals were transported on 107 seawater ice in air to the research station of the Ministère de l'Agriculture, des Pêcheries et de 108 l'Alimentation du Québec (MAPAQ) at Grande-Rivière (CAMGR), less than 50 km from the sampling site. Upon arrival at CAMGR, animals were tagged with Bee Tags<sup>™</sup> (labels were 109 110 glued on the upper valve using a cyanoacrylate adhesive) and placed in three flow-through seawater tanks (100 L) kept at the water temperature measured during collection (7.2°C in 111 July 5<sup>th</sup>, 7.1°C in September 6<sup>th</sup> and 6.3°C in October 18<sup>th</sup>) and held under natural 112 113 photoperiod. Seawater was filtered (1 µm) and UV-sterilized. Salinity varied between 28.2 114 and 29.1 %, as is encountered in the scallops' natural habitat. According to the stage of the 115 experiment, the scallops were either starved or fed with phytoplankton cultures (see below).

At the end of each series of metabolic and behavioural analyses at CAMGR, scallops were flown live on frozen seawater (travel time less than 12 h) to Université Laval, Québec for analysis of mitochondrial performance, dissection and gonad sampling. After their arrival, scallops were placed in a 1000 L aquarium containing artificial seawater at 7.0±0.5 °C. After 4 h of acclimation and during the subsequent 4-5 days, scallops were sampled to assess properties of muscle mitochondria, take gonad samples and measure tissue-wet masses.

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# Determination of reproductive status

124 For each individual, part of the gonad was rinsed with filtered seawater and then fixed 125 in a 10% Helly fixative (Shaw and Battle 1957). Tissues were dehydrated through an 126 ascending ethanol series and embedded in paraffin wax. Four 5µm thick sections were cut 127 through the entire piece of gonad and then stained with Ehrlich's haematoxylin and eosin 128 (H&E). Reproductive condition was assessed from one field per section (four fields per 129 scallop) using stereological analysis. For females, the proportions of normal and atresic 130 oocytes per field (gamete volume fraction, GVF, and atresic volume fraction, AVF) were 131 estimated on histological slides examined at 400x magnification with an image capture kit 132 CoolSnap-Pro digital kit 4.1 (Pernet et al. 2003). Oocytes with atretic (abnormal shape or 133 colour) or lytic (denaturated) characteristics were considered to be in resorption (Lubet 1959). 134 For males and females, the gametogenic stage (indifferent, development, mature, spawning or 135 spent) of each individual was determined by microscopic observation of slides as described 136 by Lowe et al. 1982. We also estimated the gonadosomatic index (GSI: gonad wet mass/total 137 wet mass of soft tissues) as a quantitative criterion of reproductive stage.

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#### Evaluation of escape responses

Individual scallops were placed in 60 x 60 x 12 cm trays containing ~ 15 L of filtered (1  $\mu$ m) seawater at the habitat temperature and escape responses were measured following Brokordt et al. (2000a) and Lafrance et al. (2003). After 2 min in the trays and when they were ventilating normally, scallops were stimulated to swim and escape by touching them with an arm of a sea star, *Asterias vulgaris*. The time before the first reaction of the scallop, the maximal number of valve adductions (claps) in a series, the total number of claps before exhaustion, and the escape time (total time until exhaustion minus the time before the scallop's first reaction) were noted. The observation was stopped if no claps occurred after 2
min of stimulation. After 30 min of recuperation (during which metabolic rate was measured),
each scallop underwent a second escape response test (again followed by 30 min of metabolic
rate measurement).

151 Sea stars were obtained from the same site as the scallops and were maintained in a 152 flow-through seawater tank (100 L) at temperatures similar to those of the scallop tanks. To 153 standardize their hunger level, sea stars were starved for 24 h before the experiments.

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#### Metabolic rate measurements

156 VO<sub>2</sub>min, or standard metabolic rate (SMR), estimates maintenance requirements of 157 resting, unstressed organisms that are not digesting food and are at a stable temperature within 158 their optimal range (Rolfe and Brown 1997). After being acclimated to maintenance 159 conditions and starved for three days, the oxygen consumption of each scallop was measured 160 at habitat temperature ( $\approx 7^{\circ}$ C) to estimate SMR. Animals were transferred individually into 1 161 L metabolic chambers that were maintained open for 60 min before starting measurements. 162 Four chambers were used simultaneously, which allowed us to measure three animals and a 163 blank with an empty shell. The oxygen consumption of individual animals was determined by 164 sealing the chamber and measuring the reduction in percent dissolved  $O_2$  with a YSI (5331) 165 polarographic electrode and analyzer. Seawater in the metabolic chamber was well mixed 166 with a magnetic stirrer. The output signal was monitored continuously starting from 100% 167 saturation and until at least a 20% decrease in saturation was reached. Immediately after 168 measurements, scallops were re-introduced into the flow-through seawater tanks and fed a 1:1 169 mixture of *Chaetoceros muelleri* and *Isochrysis galbana*. The diet was supplied continuously, maintaining a concentration of  $\sim 30$  cells. $\mu$ L<sup>-1</sup> in the tanks, allowing the scallops to feed to 170 171 satiety for two days (Pernet et al. 2003; Pernet et al. 2005).

172 Maximal metabolic rates (VO<sub>2</sub>max) were assessed using scallops immediately after 173 the escape response tests. Scallops were transferred into metabolic chambers and oxygen 174 consumption was measured for 30 min. Scallops were open and ventilating after a few 175 minutes in the chambers thus ensuring that animals were well into their aerobic recovery 176 phase. Therefore, their oxygen consumption most likely reflected the maximum  $O_2$ 177 consumption rate (Tremblay et al. 2006). The escape response was then quantified a second 178 time, followed by a second determination of oxygen consumption rates. The higher of the two 179 oxygen uptake rates following fatigue was considered as the maximum metabolic rate  $(VO_2max)$  (Tremblay et al. 2006). Oxygen consumption (ml O<sub>2</sub>. h<sup>-1</sup>) was expressed as the rate 180 181 expected for a standard animal (soft tissue wet mass of 50 g) using the allometric correction, v=aM<sup>b</sup>, where y is the oxygen consumption and b the scaling coefficient (Davies and Moyes 182 183 2007). We calculated a scaling coefficient of 0.8 using our data. We used wet instead of dry 184 mass as the denominator for oxygen uptake rates as the use of the muscle for subsequent 185 biochemical measurements prevented assessment of dry mass.

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# Mitochondrial isolation and measurement of substrate oxidation

188 Isolation procedures and measurements of substrate oxidation followed Guderley et al. 189 (1995) and Brokordt et al. (2000a). For mitochondrial isolation, all manipulations were 190 carried out on ice except the centrifugations, which were performed at 4°C. Phasic muscles 191 were rinsed in 5 mL isolation medium and then minced. The minced muscle was then 192 homogenized in 8 volumes of ice-cold isolation buffer containing 480 mM sucrose, 30 mM 193 HEPES, 230 mM KCl, 3mM Na<sub>2</sub>EDTA, 6 mM EGTA, and 5 mM MgCl<sub>2</sub>, pH 7.0 at 25°C. On 194 the day of the experiment, 0.1% fatty acid-free bovine serum albumin (BSA) was added. The 195 homogenate was centrifuged at 900g for 10min and the superficial lipid layer was removed. 196 The remaining supernatant was again centrifuged at 900g for 10min. The resulting 197 supernatant was considered free of unbroken cells or cell debris and was centrifuged at 9000g. 198 To optimize the purity of mitochondrial pellets and to remove MgCl<sub>2</sub>, the resulting pellet was 199 rinsed once by resuspension in isolation buffer free of MgCl<sub>2</sub> and recentrifuged at 9000g. The 200 mitochondrial pellet was re-suspended in a volume of reaction buffer corresponding to one-201 tenth of the mass of muscle used.

202 Mitochondrial oxygen uptake was measured polarographically using a water-jacketed 203 O<sub>2</sub> monitoring system (Qubit System, Kingston, Ontario, Canada). Temperature was 204 maintained at 5°C by a circulating refrigerated water bath. For each assay, around 1 mg of 205 mitochondrial protein (~100 µL mitochondrial preparation) was added to 1 mL assay medium 206 containing 480 mM sucrose, 70 mM HEPES, 100 mM KCl, 10 mM KH<sub>2</sub>PO<sub>4</sub>, 50 mM taurine 207 and  $\beta$ -alanine, pH 7.0. On the day of the experiment, 0.5% BSA was added to the assay 208 medium (Guderley et al. 1995). To obtain maximal rates of respiration (state 3), glutamate was added to a final concentration of 24 mmol. $L^{-1}$  and ADP to a final concentration of 0.4 209 210 mmol.L<sup>-1</sup>. The ADP/O ratio was measured according to Chance and Williams (1956). Only 211 mitochondrial preparations with respiratory control ratios (RCR; state 3 / state 4, when ADP 212 was depleted)  $\geq$  3 were used. Preliminary experiments established that glutamate was 213 oxidized at higher rates than pyruvate or succinate.

214

215 *Cytoc* 

# Cytochrome C oxidase activity

216 CCO activity was measured at 5°C and 15°C according to Kraffe et al. (2007). Fresh 217 mitochondrial suspensions were diluted in phosphate buffer without Triton-X (45 mmol.L<sup>-1</sup> 218 KH<sub>2</sub>PO<sub>4</sub> and 30 mmol.L<sup>-1</sup>K<sub>2</sub>HPO<sub>4</sub>, pH 6.8). We used an initial cytochrome C concentration of 219 100  $\mu$ mol.L<sup>-1</sup>. All assays were run in triplicate using fresh mitochondrial preparations. 220 Activities were calculated using an extinction coefficient of 19.1 mmol.L<sup>-1</sup>.cm<sup>-1</sup> (first order 221 reaction). 222

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#### Cytochromes, ANT and protein concentrations

224 Cytochromes A, B, C and  $C_1$  in the mitochondrial preparations were quantified by 225 difference spectra (Kraffe et al. 2007). The electron transport chain components in 2% deoxycholate-dispersed mitochondria were reduced by 5 mmol.L<sup>-1</sup> ascorbate and oxygen in 226 227 the solution was eliminated by the addition of dithionite (Williams 1964). The reduced samples were read against samples oxidized with 5 mmol.L<sup>-1</sup> ferricyanide. The concentration 228 229 of adenine nucleotide translocase (ANT) in mitochondrial suspensions was measured by 230 titration with its non-competitive irreversible inhibitor, carboxyatractyloside (CAT) (Guderley 231 et al. 2005). The quantity of ANT in mitochondrial suspensions corresponded to the amount 232 of CAT needed for inhibition. The protein concentration in mitochondrial suspensions was 233 determined by the bicinchoninic acid method (Smith et al. 1985) using BSA.

234

#### 235 Membrane lipid analysis

The membrane lipids of mitochondrial suspensions were extracted according to Folch et al. (1957) as modified for mitochondrial preparations (Kraffe et al. 2007). Separation of polar and neutral lipids on silica gel micro-columns, separation of membrane lipid classes, FA and sterol analyses, and calculation of amounts of phospholipid classes followed Kraffe et al. (2007).

# 241

# Separation of polar lipids on silica gel micro-columns

An aliquot of the lipid extract was deposited at the top of a silica gel micro-column (30 x 5 mm i.d., packed with Kieselgel 60 (70-230 mesh, Merck) previously heated at 450°C and deactivated with 6 wt%  $H_2O$  (Marty et al. 1992). Neutral lipids were eluted with 10 mL of CHCl<sub>3</sub>/MeOH (98:2, vol/vol) and stored at -20°C for later sterols analysis. The polar lipid fraction was recovered with 20 mL of MeOH and stored at -20°C for later phospholipid class

separation by high performance liquid chromatography (HPLC) and FA composition analysisby gas chromatography (GC).

#### 249 Cholesterol analysis

250 Free sterols (membrane components) and esterified sterols (sterol reservoir and 251 reserves) from the neutral lipid fraction were not previously separated since sterols are mostly 252 in the free form in scallop muscle (Napolitano and Ackman 1992; Palacios et al. 2007). An 253 aliquot of the neutral lipid fraction was transesterified with methoxide (MeONa) for 90 min at 254 ambient temperature (Soudant et al. 1996). The sterols thus released were extracted in hexane 255 and injected directly into GC. Sterols were analyzed in a Chrompak 9002 gas chromatograph 256 equipped with a RTX65 (65% diphenyl, 35% dimethylpolysiloxane) fused silica capillary 257 column (50 m x 0.32 mm, 0.2 pm film thickness) using an on-column injection system and 258 hydrogen as carrier gas, with a thermal gradient from 60 to 280°C. Quantification of sterols 259 was achieved by adding a known quantity of cholestane to samples.

# 260 Separation of membrane lipid classes and FA analysis

261 Separation of phospholipid classes and subclasses used two successive HPLC 262 separations with two different mobile phases. This method allowed the separate analysis of 263 plasmalogen (1-alkenyl-2-acyl-) and diacyl subclasses of phosphatidylethanolamine (PE), 264 phosphatidylcholine (PC) and phosphatidylserine (PS) in bivalves along with cardiolipin 265 (CL), phosphatidylinositol (PI), and natural lysophosphatidylcholine (LysoPC) (Kraffe et al. 266 2004). Each fraction was collected and, after transesterification (MeOH/BF<sub>3</sub>), analyzed by GC 267 for FA composition. Fatty acid methyl esters (FAME) obtained were identified and quantified 268 using both polar (CPWAX 52 CB – 50m x 0.25 mm i.d.; 0.2 µm thickness) and non-polar 269 (CP-Sil 8 CB – 25m x 0.25 mm i.d.; 0.25µm thickness) capillary columns and C23:0 FA as an 270 internal standard. FA were expressed as the molar percentage of the total FA content of each 271 class or subclass. For plasmalogen subclasses, the total percentage was adjusted to 50% to

take into account the absence of alkenyl chains of the *sn*-1 position hydrolyzed by the acidmobile phase.

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#### Calculation of amounts of phospholipid classes

The quantities of each phospholipids class and subclass of were determined from their respective FA spectrum obtained by GC. To obtain the molar content of each analyzed fraction, a correction factor was applied to their respective total FA molar contents: x1 for plasmalogen fractions and for the natural lysoPC fraction; x1/2 for PI and the diacyl fractions of PE, PC and PS, and x1/4 for the CL fraction.

280

### 281 Statistical analysis

282 Statistical comparisons were carried out with StatGraphics Plus 5.1 (Sigma Plus Inc., 283 Toulouse, France). Initial two-factor analysis of variance (ANOVA) (reproductive stage and 284 sex as factors) established that sex was not a significant factor for the measured parameters, 285 thus values for males and females were pooled. None of the variables we measured showed 286 any significant tank effect or interaction between tank and reproductive state. One-way 287 ANOVA followed by a posteriori Bonferoni multiple comparisons was used for analysis of 288 the effect of reproductive stage on behaviour, metabolic measurements and mitochondrial 289 characteristics. Differences were considered significant when P<0.05.

For lipid compositions, expressed as mol%, two-way factorial multiple analysis of variance (MANOVA) using SPSS 13.0 (Lead Technologies, Chicago, Illinois, USA) was applied with reproductive stage as the factor. Data were log+1 or 1/sqrt (x) transformed before the analysis. Where differences were detected, least-square means multiple comparison tests were used to determine which means were significantly different. Residuals were screened for normality using expected normal probability plots and homogeneity of variance was assessed with Levene's Test.

297 **Results** 

# 298 *Reproductive status*

299 The gamete volume fraction (GVF) and gonadosomatic index (GSI) both showed a significant effect of date (Figure 1, P<0.001). The highest values were observed on July 5<sup>th</sup> 300 301 and represented the mature (or ripe) gametogenic stage; lower values were observed on 302 September 6<sup>th</sup> and October 18th. Maturation stage as determined by microscopy indicated that 303 all male (2) and female (7) scallops in July had mature gonads (pre-spawned). Although GVF 304 and GSI did not discriminate between scallops in the September and October samples, 305 maturation stages did. In September, all female gonads (3) were spent while 4 of the 6 males 306 analyzed were virtually spent and 2 were spent. In October, all male gonads (3) were spent, 1 307 of the 5 female gonads was spent and 4 showed small primary oocytes. Thus, we identified 308 scallops sampled in early September as spawned and those in October as reproductive 309 quiescent.

310

#### 311 Escape response behaviour

312 All behavioural responses except for the maximum number of claps in a series, 313 changed with reproductive stage (Figure 2). Specifically, reproductive-quiescent scallops took 314 more time to respond to the predator both for their initial test (P<0.001) and after 30 min of 315 recuperation (P=0.03) (Figure 2a). Spawned scallops made fewer claps before exhaustion 316 than did pre-spawned and reproductive-quiescent scallops (P=0.03) (Figure 2c). Total escape 317 time was shorter for spawned scallops during their initial response and after 30 min of 318 recuperation (P<0.001) (Figure 2d). Spawned scallops tended to have a lower capacity for 319 recovery, only performing 56% of their initial claps after recovery whereas scallops in the 320 other reproductive stages responded with 68-75% of their initial claps (Table 1). After 30 min 321 of recuperation, the escape responses of pre-spawned and reproductive-quiescent scallops lasted 82-85% of their initial duration while spawned animals only responded for 53% of their
initial escape duration (P=0.02) (Table 1).

- 324
- 325 *Aerobic power budget*

326 Metabolic rates after exhaustion (VO<sub>2</sub>max) did not vary significantly with 327 reproductive state, but VO<sub>2</sub>min did, with higher values of oxygen consumption in spawned 328 scallops than at the other stages (P<0.001) (Figure 3). Whole animal metabolic rates and 329 allometrically corrected rates showed the same trends between reproductive stages. 330 VO<sub>2</sub>min/VO<sub>2</sub>max ratios, indicating the proportion of oxygen consumption needed for 331 maintenance, showed values of 34 and 41% in pre-spawned and reproductive-quiescent 332 animals and significantly higher values in spawned scallops (59%, P=0.005) (Figure 3). 333 Aerobic scope, i.e. the difference between maximum and standard  $VO_2$ , tended to be lowest in 334 spawned scallops (Figure 4).

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336

#### Oxidative capacities and composition of muscle mitochondria

(nmol O.min<sup>-1</sup>.mg<sup>-1</sup> 337 oxidation (State 3) Maximal rates of glutamate 338 mitochondrial protein) did not change with reproductive stage (Table 2). State 4 rates (nmol O. min<sup>-1</sup>.mg<sup>-1</sup> mitochondrial protein) and the phosphorylation capacity of mitochondria, 339 340 expressed as the molar ratio between added ADP and consumed oxygen (ADP/O), also 341 changed little between reproductive stages.

The denominator typically used to standardize mitochondrial rates is the protein content in the mitochondrial preparation. To evaluate rates of oxygen uptake relative to parameters that are exclusively located in mitochondrial membranes, we used the concentrations of ANT and cytochromes A, B and  $C_1$  as denominators. Cytochrome levels (expressed relative to the mg of proteins in the mitochondrial preparations) did not change

347 significantly between reproductive stages except that cytochrome B was higher in spawned 348 and reproductive-quiescent scallops compared to pre-spawned animals (P=0.02) (Table 3). 349 The protein-specific levels of ANT exceeded those of the cytochromes and were significantly 350 higher (P<0.001) in reproductive-quiescent scallops compared to pre-spawned and spawned 351 scallops. Modifications in the maximal rates of glutamate respiration were more apparent 352 when mitochondrial membrane proteins were used as the denominator. When State 3 rates 353 were expressed over cytochrome B, they were higher in pre-spawned scallops than in the 354 other reproductive states (P=0.07). When expressed per nmol of ANT, mitochondrial rates 355 were significantly lower in reproductive-quiescent scallops than in the other reproductive 356 stages (P=0.02) (Figure 5).

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358

Cytochrome C oxidase activity

359 CCO activity in mitochondrial suspensions was nearly two-fold higher in 360 reproductive-quiescent animals than in the other reproductive stages at both 5 and 15°C 361 (P<0.001) (Figure 6). The same pattern was found when CCO activity was expressed over 362 cytochrome A levels, except that spawned scallops showed an intermediate catalytic capacity. 363 The  $Q_{10}$  for CCO remained between 1.7 and 1.85 for the different reproductive stages. 364 Whereas all of the measured CCO activity was used by mitochondria oxidizing glutamate 365 (State 3/CCO activity expressed in mU) in pre-spawned and spawned scallops ( $100.9\% \pm 10.3$ 366 and 109.9%  $\pm$  3.5, respectively), the proportion decreased to 71.8%  $\pm$  6.4 in reproductive-367 quiescent scallops (P=0.006).

368

# 369 *Composition of mitochondrial membrane lipids with reproductive stage*

The HPLC method used in the present study allowed the separate analysis of plasmalogen (1-alkenyl-2-acyl-) and diacyl subclasses of phosphatidylethanolamine (PE),

phosphatidylcholine (PC) and phosphatidylserine (PS) found in scallops, along with phosphatidylinositol (PI), cardiolipin (CL), and natural lysophosphatidylcholine (LysoPC). The diacyl forms of phosphatidylcholine (diacylPC), phosphatidylethanolamine (diacylPE), and phosphatidylserine (diacylPS), as well as phosphatidylinositol (PI) and cardiolipin (CL) were the predominant phospholipid classes in the mitochondrial fraction from scallop muscle (Figure 7). Plasmalogen forms were found in PE (PlsmPE) while they were only present in trace amounts in PC and PS. LysoPC was also always found in trace amounts.

The relative levels (mol%) and content (nmol.mg<sup>-1</sup> protein) of glycerophospholipids 379 380 (total, classes and subclasses) remained stable between the three reproductive stages (Figure 7 381 and Table 4). Total sterol content did not change significantly with reproductive stage and 382 constituted one third of total lipids (glycerophospholipids + sterols) in the mitochondrial 383 preparations. Sterols were principally composed of cholesterol (28%), 24-384 methylenecholesterol (18%), brassicasterol (15%), 22-dehydrocholesterol (9%), 385 norcholesterol (5%) and  $\beta$ -sitosterol (5%) and the relative levels did not change between the 386 reproductive stages (data not shown).

387

388 Fatty acyl chain composition of membrane phospholipid classes and subclasses
389 between reproductive stages

The total phospholipids were dominated by 16:0 for the saturated fatty acids (SFA) and by 20:5n-3 and 22:6n-3 as the main unsaturated FA. The other major FA were 18:0 and 18:1n-7. The FA composition of total phospholipids was remarkably stable between the three reproductive stages (Table 5). Few modifications were detected in the FA compositions of the specific phospholipid classes and subclasses of mitochondrial fractions (data not shown). A major exception was the FA composition of the minor phospholipid class CL (Figure 8). The predominant FA, 22:6n-3, was higher in reproductive-quiescent scallops than in pre-

397	spawned scallops, (P=0.02) whereas 20:5n-3 represented 10% of the FA in pre-spawned and
398	spawned scallops and decreased threefold in reproductive-quiescent animals (P<0.001).
399	Levels of 18:0 were also lower in reproductive-quiescent scallops compared to pre-spawned
400	scallops (P=0.02). The decreases in 20:5n-3 and 18:0 accounted for the increase of 22:6n-3.
401	

101

403

# 404 Metabolism and behavioural responses

405 Standard metabolic rates (SMR) of *P. magellanicus* varied with reproductive stage. In 406 numerous bivalves, oxygen uptake varies seasonally in relation to ambient temperature and 407 food availability, both of which are linked with the energy demands of gametogenesis 408 (MacDonald and Thompson 1986; Shumway et al. 1988; Delgado and Pérez Camachao 2007 409 Tran et al. 2008). The influence of temperature on SMR found for scallops P. magellanicus in 410 our study cannot explain the differences in SMR since temperature varied little between 411 sampling dates and thus, during metabolic rate measurements. Standard metabolic rates were 412 higher in spawned scallops, than in pre-spawned or reproductive-quiescent scallops. The 413 lower SMR in pre-spawned scallops is not expected for bivalves during gametogenesis 414 (Shumway et al. 1988; Delgado and Pérez Camachao 2007). Nevertheless, O<sub>2</sub>-consumption in 415 the oyster C. gigas was shown to be lower in mature pre-spawned animals than after they had 416 spawned (Soletchnik et al. 1997, Tran et al. 2008). In our study, the increase in SMR after 417 spawning remains to be explained. The scallops sampled in July were likely close to 418 spawning and had probably completed gametogenesis, as reflected by oocytes maturity. At 419 Percé, in the Baie des Chaleurs, spawning takes place 2 or 3 months after the spring 420 phytoplankton bloom in May/June (Bonardelli et al. 1996). Once the gametes are mature, 421 scallops may have lower energy demands than during periods of gonad development, which 422 require high rates of food ingestion and mobilization of energy reserves. Tran et al. (2008) 423 suggested that cardiac output could be limited by mechanical pressure on the ventricle, since 424 the large gonadal volume in pre-spawning bivalves could limit blood flow and oxygen 425 consumption. In spawned scallops, the increased SMR may reflect enhanced energetic needs 426 during and after spawning, including the needs for tissue restoration.

427 While we observed a marked increase in standard metabolism of spawned scallops, no 428 significant changes in maximum metabolic rate with reproductive stage were found. 429 Consequently, VO<sub>2</sub>min/VO<sub>2</sub>max ratios (proportion of VO<sub>2</sub> required for maintenance 430 requirements) differed between reproductive stages, indicating a greater maintenance demand 431 in spawned scallops. Similarly, aerobic scope (i.e., the difference between maximum and 432 standard  $VO_2$ ) tended to be lower in spawned than in pre-spawned and reproductive-quiescent 433 scallops (Figure 4). The weakness of spawned scallops in their initial escape test as well as in 434 the capacity for recuperation may reflect this aerobic power budget. Indeed, spawned scallops 435 had a shortened total escape time and made fewer claps before exhaustion than pre-spawned 436 and reproductive-quiescent scallops. Even with this weaker initial escape response, spawned 437 scallops recovered a lower proportion of their initial status after a 30 min rest. Given the 438 patterns of metabolic support for swimming, the reduced initial escape response in spawned 439 scallops could reflect reduced arginine kinase activity or arginine phosphate levels in the 440 adductor muscle of spawned animals, such as seen in Chlamys islandica (Brokordt et al. 441 2000a) and Euvola ziczac (Brokordt et al. 2000b). In Chlamys islandica, Euvola ziczac and 442 Argopecten purpuratus, gametogenesis and spawning also reduce recuperation from 443 exhausting exercise (Brokordt et al. 2000a; Brokordt et al. 2000b; Brokordt and Guderley 444 2004). Since complete recuperation of phosphoarginine levels after exhausting exercise 445 requires aerobic metabolism (Livingstone et al. 1981), the increased maintenance 446 requirements of spawned scallops would slow their metabolic recuperation.

In contrast to responses observed for *C. islandica* and *E. ziczac*, but similar to results for *Argopecten purpuratus* (Brokordt et al. 2006), escape response behaviours measured at different points in the reproductive cycle varied both in initial tests and during repeat escape tests. In *P.magellanicus* as in *A. purpuratus*, various parameters, including the time to first respond to the sea star, the number of initial claps and the initial clapping time changed with

452 reproductive investment. In A. purpuratus, the impact of reproductive investment differed 453 between domesticated and wild scallops (Brokordt et al. 2006). Both in P. magellanicus and 454 A. purpuratus, the most marked changes were in the time to first respond to the predator. It is 455 unlikely that this parameter is linked with the changed aerobic power budget. However, the 456 decreased escape performance in spawned scallops could have a metabolic basis. We suggest 457 that, for the various species of scallops for which an impact of reproductive investment upon 458 escape response performance has been noted, limitations in the aerobic power budget may be 459 a central mechanism affecting performance.

460

# 461 Mitochondrial rates of glutamate oxidation and CCO activity

462 Mitochondrial rates of glutamate oxidation (expressed over mg mitochondrial protein) 463 changed little between sampling periods, as found in other scallops (Boadas et al. 1997; 464 Brokordt et al. 2000a; Brokordt and Guderley 2004). This lack of change in maximal 465 capacities of muscle mitochondria for glutamate parallels the lack of change of organismal 466 maximal aerobic capacities ( $VO_2max$ ). The limited changes in the capacities for glutamate 467 oxidation of muscle mitochondria (per mg protein) do not preclude modifications in the 468 capacity for oxidation of other substrates, such as pyruvate. Indeed, during gonadal 469 maturation and immediately after spawning, Chlamys islandica have lower mitochondrial 470 capacities for pyruvate oxidation (Brokordt et al. 2000a). Pyruvate being the principal 471 substrate to be oxidized after muscular activity, this reduced capacity for pyruvate oxidation 472 was suggested to slow the aerobic recovery of scallops from exhausting escape responses 473 (Guderley et al. 1995; Brokordt et al. 2000a).

474 Numerous enzymatic complexes participate in oxidative phosphorylation, with
475 complex IV (CCO) having significant control over mitochondrial respiration rates (Groen et
476 al. 1982; Blier and Lemieux 2001). In scallop muscle mitochondria, CCO activity did not

follow the same patterns as rates of glutamate oxidation expressed over proteins, cytochromes or ANT. This suggests that in scallop muscle, maximal rates of glutamate respiration (state 3 rates) are not dictated, solely or in part, by changes in CCO capacity. The calculated proportion of maximal CCO capacity used by the mitochondria oxidizing glutamate (State 3/CCO activity expressed in mU) was lower in reproductive-quiescent scallops than in pre- or spawned scallops. These calculated values are high compared to those for fish red muscle (Blier and Lemieux 2001; Kraffe et al. 2007) and may be due to differing assay conditions.

484

#### 485 Membrane lipid composition

486 Proportions of phospholipid classes and subclasses as well as those of sterols did not 487 change with reproductive state. Few modifications were apparent for the FA composition of 488 total phospholipids or of specific phospholipid classes and subclasses. However, among the 489 isolated classes, marked differences were found in the FA composition of CL. This minor 490 phospholipid class, predominantly constituted of 22:6n-3 in scallops (Kraffe et al. 2002), 491 showed fairly high proportions (10 mol% of the total FA in CL) of 20:5n-3 in pre-spawned 492 and spawned scallops whereas this FA in reproductive-quiescent scallops was three fold less 493 concentrated and 22:6n-3 was present at a significantly higher level. Interestingly, the 494 decrease in 20:5n-3 (and the increase in 22:6n-3) in CL coincided with the increase in CCO 495 activity.

496 Control of membrane-bound protein complexes in mitochondria (CCO, cytochrome 497  $bc_1$  complex, ADP-ATP translocase) is exerted partly at the level of CL (Schlame et al. 2000; 498 Paradies et al. 2002; Schlame and Ren 2006), with changes in the FA composition of CL 499 influencing, at least in part, the activity of these membrane complexes. In particular, 500 alterations in the CL acyl composition are suggested to modulate the CCO activity in 501 mammals (Yamaoka et al. 1988; Robinson et al. 1990; Berger et al. 1993; Watkins et al.

502 1998) as well as in fish (Wodtke 1981; Kraffe et al. 2007). We found that changes in CCO 503 activity in scallop muscle mitochondria with reproductive stage paralleled FA modifications 504 of this annular phospholipid. We propose that the increase of CCO activity in reproductive-505 quiescent scallops is due, at least in part, to decreased levels of 20:5n-3 in CL concomitant 506 with increases in 22:6n3.

507

#### 508 Standard metabolic rate and muscle mitochondrial properties

509 Basal mitochondrial proton conductance (or proton leak) may account for 20-25% of 510 SMR (Rolfe et al. 1999). Although quantitative estimates of the contribution of mitochondrial 511 proton leak to standard metabolic rate are not available for scallops, the mitochondrial carrier 512 protein, ANT, and the FA composition of inner-membrane phospholipids can explain much of 513 the variation in basal proton conductance in many animals (Brand et al. 2005; Hulbert and 514 Else 2005). If ANT were an important determinant of basal proton conductance in scallop 515 muscle mitochondria, then mitochondria in reproductive quiescent scallops, with their higher 516 ANT contents, would have a higher proton leak. However, the lack of change of state 4 rates 517 with reproductive stage suggests that proton leak does not increase with ANT levels. 518 Furthermore, the minor changes in the FA composition of total phospholipids in muscle 519 mitochondria suggest that they would contribute little to shifts in proton conductance between 520 reproductive stages. Thus, the increased SMR in spawned scallops is more likely associated 521 with other energetic needs or mechanical routes.

522

#### 523 Conclusion

524 This study examined mechanisms that could explain changes in escape response 525 performance of giant scallop, *Placopecten magellanicus*, with reproductive status. We found a 526 rise in standard metabolic rate without significant changes in maximum metabolic rate in 527 spawned scallops, suggesting that increased maintenance requirements and decreased aerobic 528 scope limit their capacity to escape and to recover from exhaustion. While examining muscle 529 mitochondrial properties, we found that glutamate oxidation capacity did not change with 530 reproductive status, although CCO did. Even though our examination at these multiple levels 531 did not seem to draw a single causal chain between the molecular and organismal levels, links 532 were apparent between some levels of organization. The stability of maximal organismal  $VO_2$ 533 paralleled the lack of change of maximal capacities for glutamate oxidation by phasic muscle 534 mitochondria. This suggests a possible role of muscle mitochondria in setting organismal 535 VO<sub>2</sub>max. CCO activity modification was associated with changes in the FA composition of 536 CL. In keeping with the regulatory role of CL for major complexes in oxidative 537 phosphorylation, the marked changes in 20:5n-3 and 22:6n-3 of CL suggest that the specific 538 FA composition of CL may modify CCO activity in scallop muscle.

539

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546

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- 705 FIGURE LEGENDS
- 706

**Figure 1:** Gamete volume fraction (GVF) and gonadosomatic index (GSI; gonad wet mass/ total tissue mass) of scallops sampled at different reproductive stages. GSI was determined for males and females while GVF was only determined for females. Values are means ± s.e.m. (GVF: N=7 for pre-spawned scallops, N=3 for spawned and N=5 for reproductive-quiescent scallops; GSI: N=8 for pre-spawned scallops, N=7 for spawned and reproductive-quiescent scallops). Different letters indicate values that differ between dates and hence reproductive stages (ANOVA and *a posteriori* test, P<0.05).

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Figure 2: Escape responses performance as a function of reproductive status of *Placopecten magellanicus*. A) Time before first reaction, B) maximum number of claps in a series, C) total number of claps and D) total escape time before exhaustion for scallops at the different reproductive stages. Values are means  $\pm$  s.e.m. (N=9 for pre-spawned and spawned scallops, N=8 for reproductive-quiescent scallops). Different letters indicate values that differ between reproductive stages (ANOVA and *a posteriori* test, P<0.05).

721

Figure 3: Standard metabolic rates (VO<sub>2</sub>min) and metabolic rates after exhaustion (VO<sub>2</sub>max) for scallops at different reproductive stages. Values are means  $\pm$  s.e.m. (N=9 for pre-spawned and spawned scallops, N=8 for reproductive-quiescent scallops). Different letters indicate values that differ between reproductive stages (ANOVA and *a posteriori* test, P<0.05).

726

Figure 4: Ratio of standard metabolic rate to maximal VO<sub>2</sub> and maximum minus standard metabolic rates (VO<sub>2</sub>max - VO<sub>2</sub>min) for scallops at different reproductive stages. Values are means  $\pm$  s.e.m. (N=9 for pre-spawned and spawned scallops, N=8 for reproductive-quiescent scallops). Different letters indicate values that differ between reproductive stages (ANOVA and *a posteriori* test, P<0.05).

732 733

**Figure 5**: State 3 respiratory rates (nmol  $O.min^{-1}$ ) expressed over concentrations of cytochromes and ANT (nmol<sup>-1</sup>) in mitochondria isolated from the muscle of scallops at different reproductive stages (assay temperature, 5°C). Values are means  $\pm$  s.e.m. (N=8 for pre-spawned scallops, N=7 for spawned and reproductive-quiescent scallops). Different letters indicate rates that differ between reproductive stages (ANOVA and *a posteriori* test,
P<0.05).</li>

740

Figure 6: Cytochrome *C* oxidase (CCO) activity in muscle mitochondria isolated from scallops at different reproductive stages, expressed in  $U_{CCO}$  mg<sup>-1</sup> mitochondrial protein and  $U_{CCO}$  nmol<sup>-1</sup> cytochrome A, measured at 5°C (white bars) and 15°C (grey bars) ( $U_{CCO}=\mu$ mol cytochrome *c* reduced min<sup>-1</sup>). Values are means ± s.e.m. (N=8 for pre-spawned and spawned scallops and N=7 for reproductive-quiescent scallops). Different letters indicate rates that differ between reproductive stages at a given assay temperature (ANOVA and *a posteriori* test, P<0.05).

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**Figure 7**: Classes and subclasses of phospholipids from the mitochondrial fraction isolated from scallop muscle at different reproductive stages. Phospholipids are expressed as mol% of total moles of glycerophospholipids. Values are means  $\pm$  s.e.m. (N=4 for each stage). No significant differences were noted between reproductive stages (two-way factorial MANOVA, least square means multiple comparison tests, P<0.05).

\*diacyl form of phosphatidylserine (PS), phosphatidylethanolamine (PE) andphosphatidylcholine (PC).

\*\*plasmalogen (1-alkenyl-2-acyl-) form of phosphatidylethanolamine (PE). The plasmalogen
forms of PC and PS were also detectable but only in trace amounts (< 0.5mol%).</li>

758

**Figure 8**: Main fatty acid composition of cardiolipin (CL) in mitochondrial fractions isolated from scallop muscle at different reproductive stages. Values are means  $\pm$  s.e.m. (N=4 for each stage). Different letters indicate values that differ between reproductive stages (two-way factorial MANOVA, least square means multiple comparison tests, P<0.05).

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**Table 1:** Recuperation (%) of total claps and escape time after 30 min following exhausting765exercise for scallops in the different reproductive states. Values are means  $\pm$  s.e.m. (N=9 for766pre-spawned and spawned scallops, N=8 for reproductive-quiescent scallops). Different767superscripts indicate differences between reproductive stages (ANOVA and *a posteriori* test,768P<0.05).</td>

% Recovery	Pre-spawned	Spawned	Reproductive quiescent
Total claps	$75.2 \pm 9.8$	$56.2 \pm 7.6$	$67.8 \pm 8.1$
Escape time	$84.5 \pm 11.1^{a}$	$53.3 \pm 7.3^{b}$	$81.7 \pm 5.1^{a}$

771 **Table 2:** Oxidative capacities at 5°C of muscle mitochondria from scallops in different 772 reproductive stages. State 3 and state 4 rates of glutamate oxidation are expressed as nmol 773  $O.min^{-1}.mg^{-1}$  mitochondrial protein. Values are means  $\pm$  s.e.m. (N=8 for pre-spawned 774 scallops, N=7 for spawned and reproductive-quiescent scallops). No significant differences 775 were noted between reproductive stages.

	Pre-spawned	Spawned	Reproductive quiescent
State 3	$46.3 \pm 4.6$	$53.5 \pm 3.8$	$53.0 \pm 3.6$
State 4	8.9 ± 1.3	$7.7 \pm 0.7$	$9.1 \pm 0.6$
ADP/O	$2.5 \pm 0.2$	$2.6 \pm 0.1$	$2.3 \pm 0.1$

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**Table 3:** Concentration of cytochromes and ANT in scallop muscle mitochondria at different778reproductive stages. Cytochrome and ANT concentrations were normalized to the protein779content in the mitochondrial preparations. Values are means  $\pm$  s.e.m. (N=9 for pre- and780spawned scallops, N=8 for reproductive-quiescent scallops). Different superscripts indicate781values that differ between reproductive stages (ANOVA and *a posteriori* test, P<0.05).</td>

	Pre-spawned	Spawned	Reproductive quiescent
А	$0.23 \pm 0.03$	$0.18 \pm 0.01$	$0.22 \pm 0.03$
В	$0.1 \pm 0.01^{a}$	$0.16 \pm 0.02$ <sup>b</sup>	$0.15\pm0.01$ $^{\mathrm{ab}}$
<b>C</b> <sub>1</sub>	$0.16 \pm 0.02$	$0.11 \pm 0.01$	$0.15 \pm 0.01$
С	$0.05 \pm 0.01$	$0.05 \pm 0.01$	$0.04 \pm 0.01$
ANT	$0.66\pm0.07$ $^{\rm a}$	$0.75 \pm 0.02$ <sup>a</sup>	$1.05 \pm 0.05$ <sup>b</sup>

**Table 4**: Content of cholesterol, total phospholipids and phospholipid classes and subclasses785in mitochondrial fractions isolated from scallop muscle at different reproductive stages.786Values are means  $\pm$  s.e.m. (N = 4 for each stage). Different superscripts indicate values that787differ between reproductive stages (ANOVA and *a posteriori* test, P<0.05)</td>

	Pre-spawned	Spawned	Reproductive quiescent
Total glycerophospholipids (µmol.mg <sup>-1</sup> prot)	$0.31\pm0.03$	$0.37\pm0.02$	$0.38\pm0.08$
Sterols (µmol.mg <sup>-1</sup> prot)	$0.16\pm0.03$	$0.18\pm0.02$	$0.19 \pm 0.01$
% Sterols	$34.1 \pm 1.3$	$33.1 \pm 2.8$	$31.8\pm4.1$
Classes and subclasses (nmol.mg <sup>-1</sup> prot)			
CL	$1.7 \pm 0.2$	$2.1 \pm 0.2$	$1.9 \pm 0.3$
PS	$12.9\pm0.9$	$15.9\pm1.2$	$14.2 \pm 3.4$
DiacylPE	$34.8 \pm 2.9$	$35.9 \pm 2.2$	$39.4\pm8.7$
PlsmPE	$15.9 \pm 3.5$	$16.4 \pm 1.0$	$19.7 \pm 3.6$
DiacylPC	$90.9 \pm 12.9$	$114.8 \pm 7.5$	$104.9\pm22.9$
PI	$17.0 \pm 1.0^{a}$	$7.9 \pm 1.1^{b}$	$9.5 \pm 2.7^{b}$

790**Table 5**: FA composition of total phospholipids in mitochondrial fractions isolated from791scallop muscle at the different reproductive stages. Results are expressed as mol%. Values are792means  $\pm$  s.e.m. (N=4 for each stage). Different superscripts indicate values that differ between793reproductive stages (two-way factorial MANOVA, least square means multiple comparisons,794P<0.05).</td>

\*Others : Total of 19 detectable fatty acids (iso17:0, ant17:0, 15:0, 17:0, 16:1n-5, 18:1n-5, 18:2n-4, 18:3n-6,

796 18:3n-3, 20:2nmi(5,11), 20:2nmi(5,13), 20:3nmi(5,11,14), 20:2n-6, 20:3n-6, 20:4n-3, 21:5n-3, 22:2nmi(7,13),

797 22:2nmi(7,15), 22:3nmi(7,13,16), 22:4n-6), none of which exceeded 1.0%.

798 \*\*Total dimetylacetals (mainly 16:0DMA, 18:0DMA and 20:1DMA).

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Total Phospholipids			
Pre-Spawned	Spawned	Reproductive quiescent	
$2.0 \pm 0.4$	$2.3 \pm 0.2$	$2.0 \pm 0.1$	
$18.7 \pm 0.5$	$19.6 \pm 0.2$	$19.1 \pm 0.9$	
$4.9\pm0.2^{\rm a}$	$4.6 \pm 0.1^{a}$	$6.2 \pm 0.5^{b}$	
$2.8 \pm 0.3$	$3.2 \pm 0.2$	$2.6 \pm 0.2$	
$1.8 \pm 0.1$	$1.8 \pm 0.1$	$1.8 \pm 0.1$	
$5.7 \pm 0.2$	$6.0 \pm 0.2$	$5.5 \pm 0.2$	
$0.4 \pm 0.1^{a}$	$0.5 \pm 0.1^{ab}$	$0.6 \pm 0.1^{b}$	
$0.5 \pm 0.05$	$0.5 \pm 0.02$	$0.6 \pm 0.02$	
$0.6 \pm 0.1$	$0.6\pm0.04$	$0.6 \pm 0.02$	
$0.6 \pm 0.03$	$0.9 \pm 0.1$	$0.9 \pm 0.04$	
$3.0 \pm 0.2^{a}$	$2.0\pm0.2^{b}$	$2.0 \pm 0.2^{b}$	
$1.0 \pm 0.1^{a}$	$1.2 \pm 0.1^{ab}$	$1.4 \pm 0.1^{b}$	
$24.5 \pm 0.7$	$22.8\pm0.7$	$22.7 \pm 0.4$	
$0.3 \pm 0.01^{a}$	$0.4 \pm 0.01^{b}$	$0.4 \pm 0.02^{b}$	
$0.9 \pm 0.1$	$0.8 \pm 0.02$	$0.8 \pm 0.05$	
$21.3 \pm 0.3$	$21.8 \pm 0.3$	$21.3 \pm 0.6$	
$6.4 \pm 0.7$	$6.4 \pm 0.2$	$5.5 \pm 0.2$	
$4.6\pm0.7$	$4.3 \pm 0.5$	$5.0 \pm 0.2$	
$27.8 \pm 1.1$	$29.2 \pm 0.3$	$30.3 \pm 1.6$	
$13.4 \pm 0.4$	$13.9 \pm 0.3$	$12.5 \pm 0.6$	
$54.2 \pm 1.0$	$52.7 \pm 0.5$	$52.2 \pm 1.1$	
$295.9 \pm 5.0$	$288.9 \pm 2.9$	$284.2 \pm 6.4$	
	$\begin{tabular}{ c c c c c } \hline Pre-Spawned \\ \hline $2.0 \pm 0.4$ \\ $18.7 \pm 0.5$ \\ $4.9 \pm 0.2^a$ \\ $2.8 \pm 0.3$ \\ $1.8 \pm 0.1$ \\ $5.7 \pm 0.2$ \\ $0.4 \pm 0.1^a$ \\ $0.5 \pm 0.05$ \\ $0.6 \pm 0.1$ \\ \hline $0.6 \pm 0.03$ \\ $3.0 \pm 0.2^a$ \\ $1.0 \pm 0.1^a$ \\ $24.5 \pm 0.7$ \\ $0.3 \pm 0.01^a$ \\ $0.9 \pm 0.1$ \\ $21.3 \pm 0.3$ \\ $6.4 \pm 0.7$ \\ $4.6 \pm 0.7$ \\ $4.6 \pm 0.7$ \\ $27.8 \pm 1.1$ \\ $13.4 \pm 0.4$ \\ $54.2 \pm 1.0$ \\ $295.9 \pm 5.0$ \\ \end{tabular}$	$\begin{tabular}{ c c c c c } \hline Total Phospholipid\\ \hline Pre-Spawned & Spawned\\ \hline \hline Pre-Spawned & 19.6 \pm 0.2\\ 18.7 \pm 0.5 & 19.6 \pm 0.2\\ 18.7 \pm 0.5 & 19.6 \pm 0.2\\ 4.9 \pm 0.2^a & 4.6 \pm 0.1^a\\ 2.8 \pm 0.3 & 3.2 \pm 0.2\\ 1.8 \pm 0.1 & 1.8 \pm 0.1\\ 5.7 \pm 0.2 & 6.0 \pm 0.2\\ 0.4 \pm 0.1^a & 0.5 \pm 0.1^{ab}\\ 0.5 \pm 0.05 & 0.5 \pm 0.02\\ 0.6 \pm 0.1 & 0.6 \pm 0.04\\ \hline 0.6 \pm 0.03 & 0.9 \pm 0.1\\ 3.0 \pm 0.2^a & 2.0 \pm 0.2^b\\ 1.0 \pm 0.1^a & 1.2 \pm 0.1^{ab}\\ 24.5 \pm 0.7 & 22.8 \pm 0.7\\ 0.3 \pm 0.01^a & 0.4 \pm 0.01^b\\ 0.9 \pm 0.1 & 0.8 \pm 0.02\\ 21.3 \pm 0.3 & 21.8 \pm 0.3\\ 6.4 \pm 0.7 & 6.4 \pm 0.2\\ 4.6 \pm 0.7 & 4.3 \pm 0.5\\ 27.8 \pm 1.1 & 29.2 \pm 0.3\\ 13.4 \pm 0.4 & 13.9 \pm 0.3\\ 54.2 \pm 1.0 & 52.7 \pm 0.5\\ 295.9 \pm 5.0 & 288.9 \pm 2.9\\ \hline \end{tabular}$	